



SEQUENCE LISTING

<110> Arthur B. Raitano  
Daniel E.H. Afar  
Aya Jakobovits  
Mary Faris  
Rene S. Hubert  
Steve Chappell Mitchell  
Douglas C. Saffran

<120> NOVEL G PROTEIN-COUPLED RECEPTOR  
UP-REGULATED IN PROSTATE CANCER AND USES THEREOF

<130> 511582002410

<140> US 10/017,066

<141> 2001-12-14

<150> US 09/680,728

<150> 2000-10-05

<150> 60/157,902

<151> 1999-10-05

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<213> Homo Sapiens

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Phe Ile Leu Ile Gly Leu Pro Gly Leu Glu Glu Ala Gln Phe Trp Leu	
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gcc ttc cca ttg tgc tcc ctc tac ctt att gct gtg cta ggt aac ttg	267
Ala Phe Pro Leu Cys Ser Leu Tyr Leu Ile Ala Val Leu Gly Asn Leu	
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Thr Ile Ile Tyr Ile Val Arg Thr Glu His Ser Leu His Glu Pro Met	
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Tyr Ile Phe Leu Cys Met Leu Ser Gly Ile Asp Ile Leu Ile Ser Thr	
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tca tcc atg ccc aaa atg ctg gcc atc ttc tgg ttc aat tcc act acc	411
Ser Ser Met Pro Lys Met Leu Ala Ile Phe Trp Phe Asn Ser Thr Thr	
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atc cag ttt gat gct tgt ctg cta cag att ttt gcc atc cac tcc tta	459
Ile Gln Phe Asp Ala Cys Leu Leu Gln Ile Phe Ala Ile His Ser Leu	
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tct ggc atg gaa tcc aca gtg ctg ctg gcc atg gct ttt gac cgc tat	507
Ser Gly Met Glu Ser Thr Val Leu Leu Ala Met Ala Phe Asp Arg Tyr	
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Val Ala Ile Cys His Pro Leu Arg His Ala Thr Val Leu Thr Leu Pro	
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Arg Val Thr Lys Ile Gly Val Ala Ala Val Val Arg Gly Ala Ala Leu	
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Ile Ser Ala Ile Gly Leu Asp Ser Leu Leu Ile Ser Phe Ser Tyr Leu	
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Leu Ile Leu Lys Thr Val Leu Gly Leu Thr Arg Glu Ala Gln Ala Lys	
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gca ttt ggc act tgc gtc tct cat gtg tgt gct gtg ttc ata ttc tat	891
Ala Phe Gly Thr Cys Val Ser His Val Cys Ala Val Phe Ile Phe Tyr	
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Val Pro Phe Ile Gly Leu Ser Met Val His Arg Phe Ser Lys Arg Arg	
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gac tct ccg ctg ccc gtc atc ttg gcc aat atc tat ctg ctg gtt cct	987
Asp Ser Pro Leu Pro Val Ile Leu Ala Asn Ile Tyr Leu Leu Val Pro	
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 35 40 45  
 Tyr Ile Val Arg Thr Glu His Ser Leu His Glu Pro Met Tyr Ile Phe  
 50 55 60

Leu	Cys	Met	Leu	Ser	Gly	Ile	Asp	Ile	Leu	Ile	Ser	Thr	Ser	Ser	Met
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Pro	Lys	Met	Leu	Ala	Ile	Phe	Trp	Phe	Asn	Ser	Thr	Thr	Ile	Gln	Phe
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Asp	Ala	Cys	Leu	Leu	Gln	Ile	Phe	Ala	Ile	His	Ser	Leu	Ser	Gly	Met
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Glu	Ser	Thr	Val	Leu	Leu	Ala	Met	Ala	Phe	Asp	Arg	Tyr	Val	Ala	Ile
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Cys	His	Pro	Leu	Arg	His	Ala	Thr	Val	Leu	Thr	Leu	Pro	Arg	Val	Thr
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Lys	Ile	Gly	Val	Ala	Ala	Val	Val	Arg	Gly	Ala	Ala	Leu	Met	Ala	Pro
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Leu	Pro	Val	Phe	Ile	Lys	Gln	Leu	Pro	Phe	Cys	Arg	Ser	Asn	Ile	Leu
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Ser	His	Ser	Tyr	Cys	Leu	His	Gln	Asp	Val	Met	Lys	Leu	Ala	Cys	Asp
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Ile	Gly	Leu	Asp	Ser	Leu	Leu	Ile	Ser	Phe	Ser	Tyr	Leu	Leu	Ile	Leu
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Thr	Cys	Val	Ser	His	Val	Cys	Ala	Val	Phe	Ile	Phe	Tyr	Val	Pro	Phe
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Ile	Gly	Leu	Ser	Met	Val	His	Arg	Phe	Ser	Lys	Arg	Arg	Asp	Ser	Pro
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Leu	Pro	Val	Ile	Leu	Ala	Asn	Ile	Tyr	Leu	Leu	Val	Pro	Pro	Val	Leu
	275						280					285			
Asn	Pro	Ile	Val	Tyr	Gly	Val	Lys	Thr	Lys	Glu	Ile	Arg	Gln	Arg	Ile
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Met	Tyr	Ala	Val	Ala	Leu	Phe	Gly	Asn	Cys	Ile	Val	Val	Phe	Ile	Val
	35						40					45			
Arg	Thr	Glu	Arg	Ser	Leu	His	Ala	Pro	Met	Tyr	Leu	Phe	Leu	Cys	Met
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Leu	Ala	Ala	Ile	Asp	Leu	Ala	Leu	Ser	Thr	Ser	Thr	Met	Pro	Lys	Ile
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Leu	Ala	Leu	Phe	Trp	Phe	Asp	Ser	Arg	Glu	Ile	Thr	Phe	Asp	Ala	Cys
			85						90					95	
Leu	Ala	Gln	Met	Phe	Phe	Ile	His	Ala	Leu	Ser	Ala	Ile	Glu	Ser	Thr
		100						105					110		
Ile	Leu	Leu	Ala	Met	Ala	Phe	Asp	Arg	Tyr	Val	Ala	Ile	Cys	His	Pro
		115					120					125			
Leu	Arg	His	Ala	Ala	Val	Leu	Asn	Asn	Thr	Val	Thr	Val	Gln	Ile	Gly
	130						135					140			

Met	Val	Ala	Leu	Val	Arg	Gly	Ser	Leu	Phe	Phe	Phe	Pro	Leu	Pro	Leu
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Leu	Ile	Lys	Arg	Leu	Ala	Phe	Cys	His	Ser	Asn	Val	Leu	Ser	His	Ser
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Tyr	Cys	Val	His	Gln	Asp	Val	Met	Lys	Leu	Ala	Tyr	Thr	Asp	Thr	Leu
			180					185					190		
Pro	Asn	Val	Val	Tyr	Gly	Leu	Thr	Ala	Ile	Leu	Leu	Val	Met	Gly	Val
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Asp	Val	Met	Phe	Ile	Ser	Leu	Ser	Tyr	Phe	Leu	Ile	Ile	Arg	Ala	Val
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Leu	Gln	Leu	Pro	Ser	Lys	Ser	Glu	Arg	Ala	Lys	Ala	Phe	Gly	Thr	Cys
225					230					235					240
Val	Ser	His	Ile	Gly	Val	Val	Leu	Ala	Phe	Tyr	Val	Pro	Leu	Ile	Gly
				245					250					255	
Leu	Ser	Val	Val	His	Arg	Phe	Gly	Asn	Ser	Leu	Asp	Pro	Ile	Val	His
			260					265					270		
Val	Leu	Met	Gly	Asp	Val	Tyr	Leu	Leu	Leu	Pro	Pro	Val	Ile	Asn	Pro
		275					280					285			
Ile	Ile	Tyr	Gly	Ala	Lys	Thr	Lys	Gln	Ile	Arg	Thr	Arg	Val	Leu	Ala
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<212> PRT

<213> Homo Sapiens

<400> 4

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			20					25					30		
Met	Tyr	Val	Val	Ala	Met	Cys	Gly	Asn	Cys	Ile	Val	Val	Phe	Ile	Val
		35					40					45			
Arg	Thr	Glu	Arg	Ser	Leu	His	Ala	Pro	Met	Tyr	Leu	Phe	Leu	Cys	Met
	50					55					60				
Leu	Ala	Ala	Ile	Asp	Leu	Ala	Leu	Ser	Thr	Ser	Thr	Met	Pro	Lys	Ile
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Leu	Ala	Leu	Phe	Trp	Phe	Asp	Ser	Arg	Glu	Ile	Ser	Ile	Glu	Ala	Cys
			85						90					95	
Leu	Thr	Gln	Met	Phe	Phe	Ile	His	Ala	Leu	Ser	Ala	Ile	Glu	Ser	Thr
		100						105					110		
Ile	Leu	Leu	Ala	Met	Ala	Phe	Asp	Arg	Tyr	Val	Ala	Ile	Cys	His	Pro
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Leu	Arg	His	Ala	Ala	Val	Leu	Asn	Asn	Thr	Val	Thr	Ala	Gln	Ile	Gly
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Leu	Ile	Lys	Arg	Leu	Ala	Phe	Cys	His	Ser	Asn	Val	Leu	Ser	His	Ser
				165					170					175	
Tyr	Cys	Val	His	Gln	Asp	Val	Met	Lys	Leu	Ala	Tyr	Ala	Asp	Thr	Leu
			180					185					190		
Pro	Asn	Val	Val	Tyr	Gly	Leu	Thr	Ala	Ile	Leu	Leu	Val	Met	Gly	Val
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Asp	Val	Met	Phe	Ile	Ser	Leu	Ser	Tyr	Phe	Leu	Ile	Ile	Arg	Thr	Val
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Leu	Gln	Leu	Pro	Ser	Lys	Sér	Glu	Arg	Ala	Lys	Ala	Phe	Gly	Thr	Cys
225					230					235					240
Val	Ser	His	Ile	Gly	Val	Val	Leu	Ala	Phe	Tyr	Val	Pro	Leu	Ile	Gly
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Leu	Ser	Val	Val	His	Arg	Phe	Gly	Asn	Ser	Leu	His	Pro	Ile	Val	Arg
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Val	Val	Met	Gly	Asp	Ile	Tyr	Leu	Leu	Leu	Pro	Pro	Val	Ile	Asn	Pro
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Ile	Ile	Tyr	Gly	Ala	Lys	Thr	Lys	Gln	Ile	Arg	Thr	Arg	Val	Leu	Ala
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aaactgggttg gggaatctcc attttttcaa tattattttc ttctttgttt tcttgctacg	180
tataattatt aatatacctga ctaggttggtg gttggagggt tattactttt cattttacca	240
tgcagtccaa atctaaactg cttctactga tggtttacag cattctgaga taagaatggg	300
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tct tat atc ttt atc ctt cag gca gtt cta caa ctc tcc tct cag gag	96
Ser Tyr Ile Phe Ile Leu Gln Ala Val Leu Gln Leu Ser Ser Gln Glu	
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gcc cgc tac aaa gca ttt ggg aca tgt gtc tct cac ata ggt gcc atc	144
Ala Arg Tyr Lys Ala Phe Gly Thr Cys Val Ser His Ile Gly Ala Ile	
35 40 45	
tta gcc ttc tac aca cct tca gtc atc tct tca gtc atg cac cgt gtg	192
Leu Ala Phe Tyr Thr Pro Ser Val Ile Ser Ser Val Met His Arg Val	
50 55 60	
gcc cgc tgt gct gtg cca cac gtc cac att ctc ctc gcc aat ttc tat	240
Ala Arg Cys Ala Val Pro His Val His Ile Leu Leu Ala Asn Phe Tyr	
65 70 75 80	

ctg ctc ttc cca ccc atg gtc aat ccc atc atc tat ggc gtt aag acc	288
Leu Leu Phe Pro Pro Met Val Asn Pro Ile Ile Tyr Gly Val Lys Thr	
85 90 95	
aag cag atc cgt gac agt ctt ggg agt att cct gag aaa gga tgt gtg	336
Lys Gln Ile Arg Asp Ser Leu Gly Ser Ile Pro Glu Lys Gly Cys Val	
100 105 110	
aat aga gag tga gga ata agt gga aaa aga gtg ggg ccc agt gaa tgc	384
Asn Arg Glu * Gly Ile Ser Gly Lys Arg Val Gly Pro Ser Glu Cys	
115 120 125	
tgt agt ggg cca ggg ctg tgc tga gag tag atg ggt cct aga ctc cac	432
Cys Ser Gly Pro Gly Leu Cys * Glu * Met Gly Pro Arg Leu His	
130 135 140	
gtt tag ttc ttt tct tgt att atg aaa aga ata aat gat gtc ctg aag	480
Val * Phe Phe Ser Cys Ile Met Lys Arg Ile Asn Asp Val Leu Lys	
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Leu Arg Lys Lys Lys Lys Lys	
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Ala Arg Tyr Lys Ala Phe Gly Thr Cys Val Ser His Ile Gly Ala Ile	
35 40 45	
Leu Ala Phe Tyr Thr Pro Ser Val Ile Ser Ser Val Met His Arg Val	
50 55 60	
Ala Arg Cys Ala Val Pro His Val His Ile Leu Leu Ala Asn Phe Tyr	
65 70 75 80	
Leu Leu Phe Pro Pro Met Val Asn Pro Ile Ile Tyr Gly Val Lys Thr	
85 90 95	
Lys Gln Ile Arg Asp Ser Leu Gly Ser Ile Pro Glu Lys Gly Cys Val	
100 105 110	
Asn Arg Glu Gly Ile Ser Gly Lys Arg Val Gly Pro Ser Glu Cys Cys	
115 120 125	
Ser Gly Pro Gly Leu Cys Glu Met Gly Pro Arg Leu His Val Phe Phe	
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